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# Carbon-Nitrogen Bond Formation Using Sodium Hexamethyldisilazide: Solvent-Dependent Reactivities and Mechanisms

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ABSTRACT: The solvent-dependent reactivity of sodium hexamethyldisilazide (NaHMDS) toward carbon-centered electrophiles reveals reactions that are poorly represented or unrepresented in the literature, including direct aminolysis of aromatic methyl esters to give carboxamides, nitriles, or amidines, depending on the choice of solvent. S<sub>N</sub>Ar substitutions of aryl halides and opening of terminal epoxides are also examined. A combination of <sup>1</sup>H and <sup>29</sup>Si nuclear magnetic resonance (NMR) spectroscopic studies using [<sup>15</sup>N]NaHMDS, kinetic studies, and computational studies reveals the complex mechanistic basis of the preferences for simple aryl carboxamides in toluene and dimethylethylamine and arylnitriles or amidines in tetrahydrofuran (THF). A prevalence of dimer- and

(Me<sub>3</sub>Si)<sub>2</sub>NNa + CO<sub>2</sub>Me NH<sub>2</sub> N CONH<sub>2</sub>

mixed dimer-based chemistry even starting from the observable NaHMDS monomer in THF solution is notable.

#### **■ INTRODUCTION**

After decades of communal disinterest in organosodium chemistry, we began trying to shine a light on its potential by expanding the toolbox for generating and using organosodium species.  $^{1-3}$  This involved some minor adjustments to protocols for generating and handling sodium diisopropylamide (NaDA)<sup>2,3</sup> as well as developing a new reagent, sodium isopropyl(trimethyl)silyl amide (NaPTA), that manifests desirable solubilities and reactivities as a strong base.<sup>4</sup> Our primary approach, however, is decidedly structural and mechanistic with the faith that understanding how solvation and aggregation influence reactivity and selectivity will propel applications through a combination of serendipity in our laboratory and need-driven progress by others. Even the most prominent organosodium reagent, sodium hexamethyldisilazide (NaHMDS), while garnering the attention of crystallographers, had evaded spectroscopic, mechanistic, and computational scrutiny until recent studies of its solvent-dependent structure and reactivity toward enolization.

This brings us to the current work. Aminolyses illustrated with generic examples in Scheme 1 are legion.  $S_N$ Ar substitutions and peptide bond formations are of unquestioned importance in pharmaceutical chemistry. With that said, any reaction in which the electrophile is merely heated in ammonia or simple alkylamine will be difficult to improve upon. However, should these simple protocols fail, owing to low reactivity or poor selectivity, the experimentalist is left with few options. In these instances, metal amides could offer solutions through the control of their coordination spheres.

We describe herein a survey of the reactivity of NaHMDS toward carbon-centered electrophiles of potential interest in

Scheme 1. Generic Aminolyses

synthesis, accompanied by detailed structural and mechanistic studies. NaHMDS plays the role of a highly reactive analogue of ammonia and a preface to expanding investigations of sodium alkylsilazides. Some of the transformations surveyed as opportunities for mechanistic studies piqued our interest as synthetically promising and have surprisingly little or no presence in the NaHMDS literature. The more mechanistically inclined will find the prevalence of dimer- and mixed aggregate-based reactivity surprising. We also inadvertently stumbled into the complex world of organosilicon chemistry.

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Table 1. Reactions of NaHMDS with Electrophiles in Various Solvents

	144-	4:4:		:-1-1	4-	1	1:4:	14	:-11
entry	substrate	conditions	product	yield	entry	substrate	conditions	product	yield
1	Ph OMe	2.0 equiv NaHMDS 25 °C, 5 h toluene	Ph NH <sub>2</sub>	72%	10	OMe 0	3.0 equiv NaHMDS 50 °C, 1 h DMEA	NH <sub>2</sub>	78%
2	OMe O	3.0 equiv NaHMDS 25 °C, 0.3 h DMEA	NH <sub>2</sub> NH <sub>2</sub>	95%	11	OMe 0 14	3.0 equiv NaHMDS/ THF 70 °C, 2.0 h 25 °C, 24 h	NH <sub>2</sub> NH 16	92%
3	OEt OEt	3.0 equiv NaHMDS 25 °C, 0.3 h toluene	$NH_2$	90%	12	17	2.0 equiv NaHMDS 25 °C, 0.05 h DMEA	NH <sub>2</sub> NH 16	95%
4	OMe O	1.0 equiv NaHMDS 50 °C, 1.0 h THF	N CN 6	86%	13	NO <sub>2</sub> F 18	2.0 equiv NaHMDS 25 °C, 2 h toluene	NO <sub>2</sub> NH <sub>2</sub> 19	85%
5	N OMe	3.0 equiv NaHMDS 50 °C, 0.3 h THF	NH <sub>2</sub>	92%	14	NO <sub>2</sub> F 20	2.0 equiv NaHMDS 25 °C, 1.0 h toluene	NO <sub>2</sub> NH <sub>2</sub>	77%
6	3 NOMe	3.0 equiv NaHMDS 70 °C, 0.3 h	7 NH <sub>2</sub>	85%	15	F NO <sub>2</sub> F 22	2.0 equiv NaHMDS 25 °C, 1.0 h toluene	NO <sub>2</sub> NH <sub>2</sub> 23	83%
7	8 OMe	3.0 equiv NaHMDS	9 N NH <sub>2</sub> NH	96%	16	N   F   24	2.0 equiv NaHMDS 110 °C, 2 h toluene	N NH <sub>2</sub> 25	55%
	8 	70 °C, 1.0 h THF	10 N		17	F N F	2.0 equiv NaHMDS 110 °C, 3 h toluene	F NH <sub>2</sub>	76%
8	OMe 0	2.0 equiv NaHMDS 70 °C, 1.0 h toluene	NH <sub>2</sub> 0	76%	18	Ph 0 28	2.0 equiv NaHMDS 60 °C, 24 h	OH Ph NH <sub>2</sub> (>50:1)	86%
9	OMe O	3.0 equiv NaHMDS 70 °C, 1.0 h THF	NH <sub>2</sub> NH <sub>2</sub> 13	95%	19	n-C <sub>6</sub> H <sub>13</sub>	toluene  2.0 equiv NaHMDS  25 °C, 24 h THF	29 OH n-C <sub>6</sub> H <sub>13</sub> NH <sub>2</sub>	76%

<sup>&</sup>lt;sup>a</sup>Forms a gel-like mixture during the reaction.

#### RESULTS AND DISCUSSION

**Aminolyses.** Substitutions of representative organic substrates by NaHMDS are summarized in Table 1. The yields are of isolated, purified products that have been desilylated during workup. <sup>1</sup>H and <sup>29</sup>Si NMR spectroscopic monitoring (vide infra) shows the reactions are often pristine and reveals unisolated silylated and sodiated intermediates. The times and temperatures listed in Table 1 are those required to achieve high conversion, as confirmed by in situ monitoring using NMR or IR spectroscopies.

For context, the aminolysis of methyl cinnamate to form carboxamide (entry 1) rather than alternatives such as 1,4-

addition, <sup>11,12</sup> enolization, <sup>5</sup> or generalized destruction caught our attention. Ester aminolyses by LiHMDS and NaHMDS may populate pharmaceutical notebooks, but they are poorly represented in the published literature. In 1963, Krüger et al. aminolyzed an aryl ester with NaHMDS to form an Omethyloximino ether, an intermediate en route to carboxamides (as in entry 2). One is detected in mechanistic studies below. <sup>13</sup> In 1998, Hwu and co-workers reported a NaHMDS-mediated conversion of aryl esters to nitriles with NaHMDS at 110–185 °C (as in entry 4), attributing a central importance to phenolic groups. <sup>14</sup> One-pot conversions of aryl esters to amidines (entries 11 and 12) or carboxamides (such as entry

2) by NaHMDS are unreported. Additions to nitriles to form amidinates (as in entry 12) are well-known for LiHMDS. Mechanistic studies below offer some thoughts on why analogous additions of NaHMDS to nitriles are rare.1

The potentially useful selectivities for the formation of carboxamides, nitriles, and amidines directly from esters have complex mechanistic underpinnings dependent on the solvent, temperature, and equivalents of NaHMDS (vide infra). The 3and 4-picoline methyl esters form gels in weak solvents (entries 9 and 11), presumably owing to head-to-tail oligomeric substrate-NaHMDS dimer complexes but without negative consequences.

S<sub>N</sub>Ar reactions (entries 15–19) represent a reaction class of unquestioned importance.8 We find it somewhat confounding that the S<sub>N</sub>Ar reactions seem to be more effective with soft nucleophiles. Malonates, for example, are far superior nucleophiles<sup>17</sup> than unstabilized enolates. We can find only one report of direct (uncatalyzed) S<sub>N</sub>Ar substitution by LiHMDS<sup>19</sup> and none for NaHMDS, while RNH<sub>2</sub>-based aminolyses facilitated by LiHMDS are legion albeit with an unknown role of the LiHMDS.<sup>20</sup> We expended considerable effort trying to glean mechanistic insights (entry 15-17) only to be thwarted by deeply colored debris that appears to be diazo derivatives arising from NaHMDS reacting with the nitro moiety noted years ago (eq 1).<sup>21</sup> Nitroarenes are spartan in the voluminous literature of lithium amide-mediated ortholithiations,<sup>22</sup> possibly for similar reasons.

A few problematic substrates are illustrated in Chart 1. Arenes 32-34 gave complex products suggestive of ortho-

Chart 1. Failed and Troublesome Substrates

metalations and possibly pyridyne intermediates. Halopyridines 35 and 36 and chloroisoguinoline 37 aminated but were prone to transfer a silyl group to the 3-position by a Fries-like process.<sup>23</sup> Nitrobenzene 38 underwent a clean halogen dance,<sup>24</sup> forming 2-fluoro-3-iodonitrobenzene.<sup>25</sup>

We replicated the highly regioselective epoxide openings reported by Withnall and co-workers in 2008<sup>26</sup> (entries 20 and 21) to evaluate them as candidates for mechanistic work and include these potentially useful results because they have gone largely unnoticed.<sup>2</sup>

To assist the reader, Scheme 2 represents a road map to forthcoming mechanistic studies. We explore the reaction coordinates to convert NaHMDS and methyl picolinate 3 to carboxamide 4 in DMEA (Part 1), to nitrile 6 in THF (Part 2),

Scheme 2. Focus of the Four Studies Examined Spectroscopically, Kinetically, and Computationally

and conversion of nitrile 6 to amidine 7 in DMEA (Part 3) and THF (Part 4). Each part delves into the underlying organosodium and organosilicon chemistry by examining spectroscopically observable structural events, kinetic studies to evaluate solvation and aggregation in the rate-limiting transition structures, <sup>28</sup> and computational probes of solvation, aggregation, critical transition structures, and elusive details along the reaction coordinates. 29,30

Monitoring the reactions by <sup>1</sup>H and <sup>29</sup>Si NMR spectroscopies showed that the reactions were very clean and provided detailed structural assignments. [15N]NaHMDS7 distinguished O-Si and <sup>15</sup>N-Si species owing to <sup>15</sup>N-<sup>29</sup>Si coupling, whereas combinations of [15N]NaHMDS and NaHMDS were exploited to examine reversible steps. DFT computations using Me<sub>3</sub>N in place of Me<sub>2</sub>NEt to reduce unnecessary degrees of freedom revealed an exothermic substitution of DMEA on dimer 39. We took the liberty of using the original numbers assigned to DMEA solvates in the schemes showing computed Me<sub>3</sub>N solvates. The shorthand of the general form  $A_mS_n$ alludes to the hexamethyldisilazide fragment (A) and solvent

Part 1: Mechanism of Aminolysis of Methyl-2-Picolinate (3) with NaHMDS/DMEA. We focused on structure—reactivity studies on additions to methyl-2-picolinate in DMEA and THF. Toluene acts similarly to DMEA but affords broader resonances. The reaction coordinate for the addition of NaHMDS to picoline 3 in DMEA observable by NMR spectroscopy is summarized in Scheme 3. Mono- and dichelated dimers 40 and 41 observable at -80 °C reflect the behavior of NaHMDS with other bifunctional ligands studied previously. Both dimers display coupling constants  $(^{1}J_{N-Si} = 8.5 \text{ and } 8.7 \text{ Hz, respectively})$  and chemical shifts (-15.8 and -16.4 ppm, respectively) characteristic of [15N]NaHMDS dimers. Addition of excess 3 affords computationally viable doubly chelated monomer 42 with an upfield <sup>29</sup>Si chemical shift and large coupling constant (-21.41 ppm,  ${}^{1}J_{N-Si} = 13.1$  Hz) characteristic of a NaHMDS monomer.<sup>7</sup> Rate studies (below) place dimer 40 on the reaction coordinate. The stability of 43 allows carboxamide 4<sup>31</sup> to be isolated in high yield (Table 1, entry 5).<sup>32</sup>

The computed structure of 40 (Figure 1) reveals a geometry that more closely approximates square planarity than tetrahedral sodium. Serial substitution from 39 to 41 is computed to be mildly exothermic for both steps (ignoring translational entropy).<sup>34</sup> DFT supports monomer stereoisomer 42, with the alternative stereoisomeric monomer (not drawn) being 10.4 kcal/mol less stable.

Scheme 3. Spectroscopically Observed Species on the Reaction of Methyl-2-Picolinate with NaHMDS/DMEA<sup>33</sup>

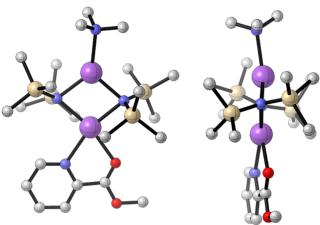
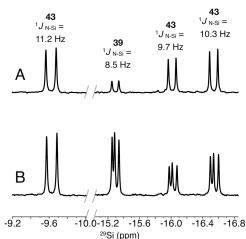


Figure 1. Ball-and-stick depictions of DFT-computed dimer 40 displaying an approximate  $10^\circ$  rotation of the chelate and  $\text{Na}_2\text{N}_2$  planes from coplanarity.

The only observable species during the aminolysis in DMEA are mixed aggregate 43 (Figure 2A) and Me<sub>3</sub>SiOMe with the latter confirmed by comparison with an authentic sample ( $^{29}$ Si,  $\delta$  17.49 ppm). Two  $^{29}$ Si resonances of 43 at -16.0 ppm ( $^{1}J_{\rm N-Si}$  = 9.7 Hz) and -16.6 ppm ( $^{1}J_{\rm N-Si}$  = 10.3 Hz) are similar to those of [ $^{15}$ N]NaHMDS dimer 39 (-15.2 ppm,  $J_{\rm N-Si}$  = 8.5 Hz) and coalesce into a single broad resonance at -20 °C. The imino  $^{29}$ Si resonance is markedly downfield (-9.70 ppm,  $J_{\rm N-Si}$  = 11.2 Hz). Addition of excess unlabeled NaHMDS to preformed [ $^{15}$ N<sub>2</sub>]43 at -80 °C shows immediate incorporation of unlabeled silazide fragments (Figure 2B). DFT computations illustrate the magnetic inequivalence of the two silazide-derived Me<sub>3</sub>Si moieties in 43 (Figure 3).

We probed the mechanism for the addition of NaHMDS to picolinate 3 in DMEA-pentane mixtures using in situ IR spectroscopy<sup>35</sup> under pseudo-first-order conditions (excess NaHMDS) following the loss of monochelated dimer 40 (1737 cm<sup>-1</sup>) to form mixed dimer 43 (1584 cm<sup>-1</sup>). Clean first-order decays (Figure 4) afford pseudo-first-order rate constants,  $k_{\text{obsd}}$ , that are independent of the initial concentration of 40. Plotting  $k_{\text{obsd}}$  vs DMEA concentration with pentane cosolvent (Figure 5) and vs NaHMDS dimer 39



**Figure 2.** Spectrum A is the <sup>29</sup>Si NMR spectrum of 0.05 M ester 3 with 0.10 M added [<sup>15</sup>N]NaHMDS in DMEA after reaction containing only mixed aggregate [<sup>15</sup>N]43 and residual NaHMDS dimer [<sup>15</sup>N]39 (Scheme 3). Spectrum B is the sample from spectrum A with 1.0 equiv (relative to ester 3) of unlabeled NaHMDS.

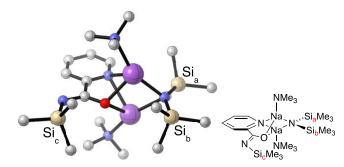
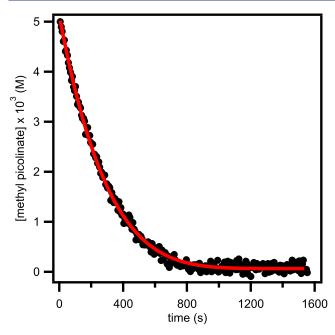
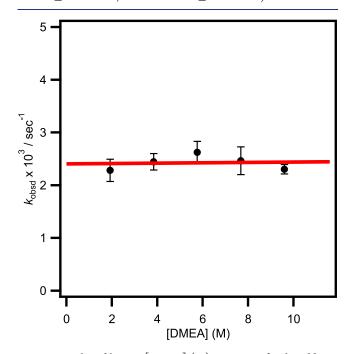


Figure 3. Ball-and-stick depictions of DFT-computed and artist's rendition of mixed aggregate 43 with Me<sub>3</sub>N as a DMEA surrogate.

concentration (Figure 6) shows zeroth-order dependencies. The resulting rate law (eq 2) is consistent with a rate-limiting addition via a transition structure of the stoichiometry  $A_2S(\text{substrate})$ . (Recall that A is an amide subunit and S is solvent.) The dimer-based reaction coordinate is a hallmark of



**Figure 4.** Addition of 0.10 M NaHMDS to methyl-2-picolinate (3, 0.005 M) at 25 °C was measured with IR spectroscopy (1737 cm<sup>-1</sup>). The curve depicts an unweighted least-squares fit to  $y = ae^{-bx}$  ( $a = 8.1 \times 10^{-3} \pm 0.1 \times 10^{-3}$ ;  $b = 1.5 \times 10^{-3} \pm 0.1 \times 10^{-3}$ ).

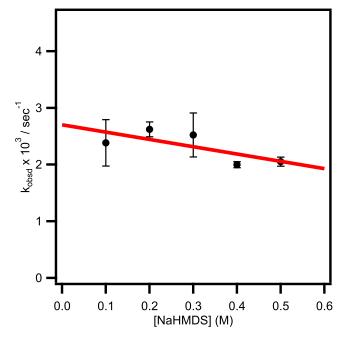


**Figure 5.** A plot of  $k_{\rm obsd}$  vs [DMEA] (M) in pentane for the addition of NaHMDS (0.10 M) to methyl-2-picolinate (3, 0.005 M) at 25 °C measured with IR spectroscopy (1737 cm<sup>-1</sup>). The curve depicts an unweighted least-squares fit to y = ax + b ( $a = 2.4 \times 10^{-3} \pm 0.2 \times 10^{-4}$ ;  $b = 3.3 \times 10^{-6} \pm 0.3 \times 10^{-6}$ ).

alkali metal amides in poorly coordinating solvents. <sup>2,7c,36</sup> Figure 7 depicts the computed transition structure **TS-1** displaying a developing Na<sub>2</sub>(O)(N) mixed dimer core.

$$-d[\mathbf{40}]/dt = k'[\mathbf{40}]^{1}[A_{2}S_{2}]^{0}[S]^{0}$$
(2)

Probing the reaction coordinate en route to observable product 43 (carboxamide 13 after workup) and Me<sub>3</sub>SiOMe



**Figure 6.** A plot of  $k_{\rm obsd}$  vs [NaHMDS] (M) for the addition of NaHMDS to methyl-2-picolinate (3, 0.050 M) in 9.6 M (neat) DMEA at 25 °C measured with IR spectroscopy (1737 cm<sup>-1</sup>). The curve depicts an unweighted least-squares fit to y = ax + b ( $a = 2.7 \times 10^{-3} \pm 0.2 \times 10^{-3}$ ;  $b = -1.3 \times 10^{-3} \pm 0.1 \times 10^{-3}$ ).

using DFT computations served up a nuanced and quite complex story (Scheme 4). To reiterate, Me<sub>3</sub>N is used as a surrogate of DMEA to reduce unnecessary degrees of freedom while taking the liberty of reusing the original numbers for 40, 43, and 46. Transition structures TS-1, TS-2, TS-3, and TS-4 all have single-negative frequencies. Intrinsic reaction coordinate (IRC) calculations<sup>37</sup> were used to determine the minima flanking each transition structure. Minima connected without connecting transition structures represent structural minima corresponding to reorganizations of insufficient interest to probe further. There are likely more minima and barriers resulting from other minor adjustments. These same caveats also apply to the reaction coordinate depicted in Scheme 6 (below).

Following the formation of open dimer 45,<sup>36</sup> the 1,2-addition proceeds via the rate-limiting transition structure TS-1 (depicted in Figure 7) to form the tetrahedral adduct as mixed aggregate 46. Structures beyond TS-1 represent kinetically and spectroscopically invisible post-rate-limiting events. The collapse of adduct 46 via transition structure TS-2 affords NaOMe–NaHMDS mixed dimer 47. Following a readjustment via TS-3 to position the methoxy as a MeO–Si Lewis acid–base complex 48, silyl transfer via transition structure TS-4 extrudes Me<sub>3</sub>SiOMe via complex 49 to form mixed dimer 50 (same structure compared to mixed aggregate 43 instead utilizing Me<sub>3</sub>N as a solvent surrogate in DFT-computation) as the spectroscopically observable product (albeit with an exothermic addition of a second solvent).

As drawn, the NaOMe silylation is an intra-aggegate reaction. We cannot exclude the possibility that NaOMe is released into solution before silyl transfer, but we have no reason to invoke it either. We also probed for a methoxy-silyl coupling directly from the tetrahedral adduct 46 without first expelling NaOMe (see 51, path a) but were unable to locate such a transfer. Similarly, a silyl transfer to extrude NaOSiMe<sub>3</sub>

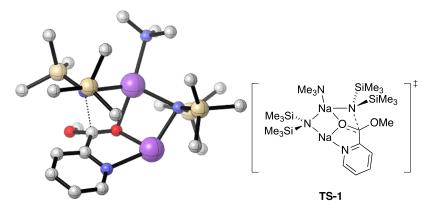
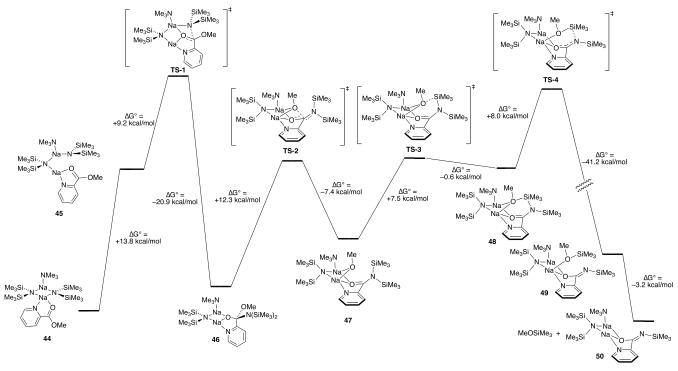
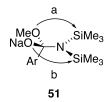


Figure 7. DFT-computed transition structure TS-1 for NaHMDS-based addition to methyl-2-picolinate (3) with Me<sub>3</sub>N as a DMEA surrogate.

## Scheme 4. DFT-Computed Reaction Coordinate for the Aminolysis of Methyl-2-Picolinate (3) by NaHMDS Using Me<sub>3</sub>N as a DMEA Surrogate



from 46 (see 51, path b) was located but had an unreasonably high (23 kcal/mol) barrier consistent with our failure to observe  $Me_3SiONa$  spectroscopically.



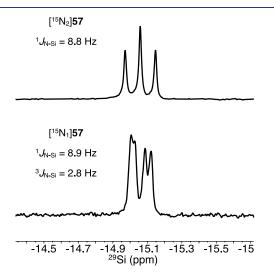
Part 2: Mechanism of Aminolysis of Methyl-2-Picolinate (3) with NaHMDS/THF. The aminolysis in THF follows a distinctly different pathway. The spectroscopically observable forms summarized in Scheme 5 show mechanistic complexity lurking beneath the surface. At the outset, we note that the excess NaHMDS undergoes a THF-dependent deaggregation of disolvated dimer 52 to provide tetrasolvated monomer 53 described previously, <sup>7a,b</sup> which proves to be mechanistically

important. Picolinate-complexed dimer 54 (analogous to 9) and THF-complexed monomer 55 can be observed at -110 °C at both low and high THF concentrations, using a pentane cosolvent. Evidence of complexation is gleaned from distinct chemical shift differences in the <sup>1</sup>H and <sup>29</sup>Si NMR spectra when compared with the uncomplexed substrate 3 and free NaHMDS (Supporting Information). The assigned THF solvation numbers derive from DFT computations.

Warming mixtures of 3 and NaHMDS to room temperature affords imino ether 56, analogous to that isolated previously, <sup>13</sup> within 1.0 min. The concomitant formation of Me<sub>3</sub>SiONa was confirmed by comparison with an authentic sample (17.70 ppm). In the absence of excess NaHMDS, 56 gives nitrile 6 over 24 h at RT or in 30 min at 50 °C in good yield (Table 1, entry 7) with concomitant formation of Me<sub>3</sub>SiOMe, confirmed with an authentic sample. The conversion of 56 to 6 is somewhat mysterious in that it does not appear to require NaHMDS and is not accelerated by excess NaHMDS, yet it does not occur by heating a crude isolated sample of 56.

Scheme 5. Spectroscopically Observed Species for the Reaction of NaHMDS to Methyl-2-Picolinate 3 in THF

Reaction of 3 with excess NaHMDS renders nitrile 6 unobservable because it is quickly scavenged to give amidinate  $57^{38}$  with noteworthy spectroscopic properties. The  $^{29}$ Si resonance of  $[^{15}N_2]57$  appears as a clean triplet owing to second-order effects referred to as "virtual coupling." The two magnetically equivalent  $^{29}$ Si nuclei experience equivalent coupling by the  $^{15}$ N nuclei despite one being proximal and one being distal to each silicon nucleus (Figure 8). Similar



**Figure 8.** <sup>29</sup>Si NMR spectra of  $[^{15}N_2]$ 57 showing virtual coupling  $(^1J_{N-Si}=8.8~{\rm Hz})$  and standard coupling in  $[^{15}N_1]$ 57  $(^1J_{N-Si}=8.9~{\rm Hz})$  and  $^3J_{N-Si}=2.8~{\rm Hz})$ .

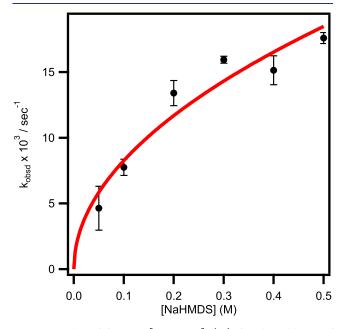
virtual coupling is observed in bridging transition metal phosphides  $(M_2P_2 \text{ core})^{39}$  and has been observed in a  $\text{Li}_2P_2$  lithium phosphide dimer. <sup>40</sup> In theory, we could have observed this in NaHMDS  $(Na_2N_2)$  dimers as well but have not. <sup>7</sup> If the symmetry is broken by generating  $[^{15}N_1]$ 57 from unlabeled nitrile 6 and  $[^{15}N]$ NaHMDS, the <sup>29</sup>Si resonance appears as a doublet of doublets owing to a large  $^1J_{N-Si}$  coupling and small  $^3J_{N-Si}$  coupling with a slight isotopic perturbation visible. <sup>41</sup>

The requisite silyl transfer from initially formed unsymmetric N,N-disilylamidine to N,N'-disilylamidine 56 is discussed in the context of DFT computations below. The

depicted 6–57 nitrile-amidinate equilibrium cannot be observed directly, but adding [<sup>14</sup>N]NaHMDS to labeled amidinate 57 results in label exchange (eq 3).

An analogous addition of NaHMDS to PhCO<sub>2</sub>Me (Table 1, entry 13) to generate phenyl-substituted amidinate 58 was approximately 60-fold (see Supporting Information) slower than for 3. It also revealed the elusive nitrile-amidinate exchange as a temperature-dependent equilibrium (eq 4). Heating excess NaHMDS (3.0 equiv) and PhCO<sub>2</sub>Me to 80 °C followed by rapid thermal quenching to -80 °C affords predominantly benzonitrile (17). Alternatively, cooling the reaction to 25 °C and letting it stand for 12 h generated amidinate 58. Heating-cooling cycles show the changes are temperature dependent and reversible. Adding [15N]NaHMDS to unlabeled benzonitrile, 17, shows label incorporation in both 17 and 58. Thus, benzonitrile can be isolated in 86% yield using 1.1 equiv of NaHMDS and elevated temperatures (Table 1, entry 13), while amidine 58 can be isolated in 98% yield using 3.0 equiv of NaHMDS at 80 °C for 3 h and then 25 °C for 12 h. This odd temperature dependence in which heating accelerates the aminolysis but retards addition to the intermediate nitrile might be why nitrile-to-amidine conversions largely exploit LiHMDS<sup>15</sup> rather than NaHMDS.<sup>16</sup> Alternatively, it could just be a cultural preference for LiHMDS. There are no extraneous silazide signals in the <sup>29</sup>Si spectrum that would implicate a mixed aggregate.

Rate studies for the addition of NaHMDS to picolinate 3 to afford imino ether **56** provided yet another nuanced story. We anticipated that the first-order dependence—the first-order decay—on complexed monomer **55** measured in neat THF would be accompanied by a zeroth-order NaHMDS dependence. To the contrary, a plot of  $k_{\rm obsd}$  vs NaHMDS shows a half-order dependence on the free NaHMDS monomer (Figure 9).

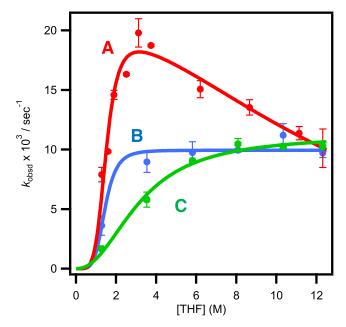


**Figure 9.** Plot of  $k_{\rm obsd}$  vs [NaHMDS] (M) for the addition of NaHMDS to methyl-2-picolinate (3, 0.005 M) in 12.8 M (neat) THF at 0 °C measured with IR spectroscopy (1731 cm<sup>-1</sup>). The curve depicts an unweighted least-squares fit to  $y = ax^n$  ( $a = 2.4 \times 10^{-3} \pm 0.1 \times 10^{-3}$ ;  $n = 0.44 \pm 0.01$ ).

The *partial* rate law (deferring discussion of solvent-concentration dependencies momentarily) is described by eq 5. To be clear on a critical point, a dimer-based addition involving contributions of a standard NaHMDS monomer would manifest a first-order NaHMDS dependence. The *half-order* on a monomer demands a full ionization of the sodium cation. 43

$$-d[55]/dt = k'[55]^{1}[AS_{4}]^{1/2}$$
  
such that  $55 = AS_{2}(3)$  (5)

We may not yet fully understand the role of solvation. The forthcoming discussion may even seem excessive, but we are still struggling to understand how organosodium solvation and aggregation differ from those of organolithiums. A plot of  $k_{\rm obsd}$ vs THF in pentane (Figure 10, curve A) manifests a maximum in the rates at intermediate THF concentrations. By contrast, analogous data using 2,5-dimethyltetrahydrofuran (2,5-Me<sub>2</sub>THF) or the decidedly more expensive 2,2,5,5-tetramethyltetrahydrofuran (2,2,5,5-Me<sub>4</sub>THF) as poorly coordinating polar cosolvents 44-47 eliminate the maximum. The solvent dependencies suggest that medium effects are in play. We remind the reader that two deaggregations are occurring concurrently. At low THF concentrations, the dimeric reactant  $A_2S$ (substrate) (54) exists concurrently with  $A_2S_2$  (NaHMDS dimer 52), and the rise in rates indicates they are collectively undersolvated relative to the rate-limiting transition structure.



**Figure 10.** A plot of  $k_{\rm obsd}$  vs [THF] (M) for the addition of NaHMDS to methyl-2-picolinate (3, 0.005 M) in 0.10 M NaHMDS at 0 °C measured with IR spectroscopy (1732 cm<sup>-1</sup>). Curve A derives from pentane cosolvent and depicts an unweighted least-squares fit to  $y = [ax^n/(1+bx^n)](1/1+cx^2)$  ( $a = 2.4 \times 10^{-3} \pm 0.1 \times 10^{-3}$ ;  $b = 1.3 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ;  $c = 0.6 \times 10^{-3} \pm 0.1 \times 10^{-4}$ ; n = 5). Curve B derives from 2,5-Me<sub>2</sub>THF cosolvent and depicts an unweighted least-squares fit to  $y = ax^n/(1+bx^n)$  ( $a = 1.6 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ;  $b = 1.6 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ; a = 5). Curve C derives from 2,2,5,5-Me<sub>4</sub>THF cosolvent and depicts an unweighted least-squares fit to  $a = 0.1 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ;  $a = 0.1 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ;  $a = 0.1 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ;  $a = 0.1 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ;  $a = 0.1 \times 10^{-1}$ ; a = 0.1

At high THF concentrations in polar cosolvents, the observable reactants  $AS_2(substrate)$  (54) and  $AS_4^{7c}$  (53) appear optimally solvated – six THF ligands total – but underaggregated relative to the dimeric rate-limiting transition structure.

The problem we confronted in this study was that 2,5-Me<sub>2</sub>THF was binding cooperatively with THF to promote monomer formation as illustrated in eq 6. At a low THF concentration, an unusually elevated concentration of NaHMDS monomer in 2,5-Me<sub>2</sub>THF compared with pentane implicates a  $(Me_3Si)_2N(THF)_x(Me_2THF)_y$  mixed solvate. However, there is almost no chance that 2,2,5,5-Me<sub>4</sub>THF cooperatively solvates monomers, and DFT computations support this assertion. We still observe the promotion of the monomer. Thus, the flattening of the curve is at least partially attributable to such an unproductive side equilibrium to form a mixed solvate according to the Principle of Detailed Balance.<sup>48</sup> We must confess that the function used to fit the THF/ pentane data (caption in Figure 10) contains a correction for inhibitory medium effects,  $1/(1+cx^2)$ , that was derived empirically with no formal molecular basis. The challenge posed by these medium effects rears its ugly head below.

With all that said, the rate law in eq 5 provides an adequate picture of a mechanism requiring 6 solvents and two silazide subunits – a hexasolvated dimer-based transition structure **TS**-5 (Figure 11) with a <sup>+</sup>Na(THF)<sub>6</sub> gegenion (Figure 12).<sup>36</sup>

We computationally examined the overall reaction coordinate using a triple-ion-based framework (Scheme 6). The cation (Figure 12) is omitted for all of the structures in Scheme 6. Triple ions are well precedented, <sup>36</sup> including for

LiHMDS. Inspection of Figure 11, however, reveals that it is a triple-ion-based reference state (59) because the lowest-energy rate-limiting transition structure TS-5 has lost all semblance of the N-Na-N triple-ion connectivity as has the IRC-derived minimum 61 preceding it.

There is plenty of room for alternative interpretation. One could, for example, imagine **61** stemming directly from a chelated monomer and ionized NaHMDS fragment. It is also possible, however, that the conversion of **60** to transition structure **TS-5** proceeds via **61** in which the  $(Me_3Si)_2N^-$  in **61** remains "bound." (The dissociation of **61** is calculated to be +5.5 kcal/mol) Transition structure **TS-5** gives way to the tetrahedral adduct **62**. The sequence follows an aza-Brook-like silicon transfer akin to that presumed to occur in an aza-Peterson-like imine formation <sup>49</sup> via transition structure **TS-6**, elimination of **63** via transition structure **TS-7** to form arene  $\pi$  complex **64**, and arene dissociation to afford the observed imino ether **56** and unobserved (fleeting) mixed triple-ion **65**.  $^{36,50}$ 

Part 3: Mechanism of Aminolysis of 2-Pyridinecarbonitrile (6) with NaHMDS/DMEA. Motivated as much by an obsessive need to complete the story as by curiosity, we examined the addition of NaHMDS to nitrile 6 to form amidinate 57 (Scheme 5) and were rewarded for our persistence. IR and NMR spectroscopy showed no evidence that nitrile 6 binds to NaHMDS at low or high DMEA concentrations. The addition occurs to give mixed amidinate 66 within seconds at 25 °C (eq 7) manifesting a silazide fragment displaying dimer-like coupling ( ${}^1J_{N-Si}=8.5$  Hz; Figure 13) that is well resolved from NaHMDS homodimer 39. As described above, further heating to 50 °C causes scrambling of the  ${}^{14}N-{}^{15}N$  isotopes.

Rate studies show an inverse-second-order dependence on DMEA (Figure 14) and first-order dependence on the NaHMDS dimer 39 (Figure 15) consistent with a disolvated-dimer-based 1,2-addition. A full computational workup is found in the Supporting Information. An

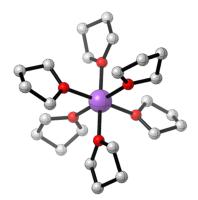


Figure 12. DFT-computed \*Na(THF)<sub>6</sub>.

abbreviated version is illustrated in Scheme 7. For example, the conversion of 70 to 66 requires several additional mundane readjustments. Also, we examined the role of solvation of the intermediates and found minima that were higher energy than the unsolvated forms except for 70, whose  $Me_3N$ -solvated form is a -2.5 kcal/mol more stable.

Part 4. Mechanism of Aminolysis of 2-Pyridinecarbonitrile (6) with NaHMDS/THF. IR and NMR spectroscopy showed no evidence that nitrile 6 binds to NaHMDS (eq 8). The only species observable by <sup>1</sup>H and <sup>29</sup>Si NMR spectroscopy was free NaHMDS (52 and 53, Scheme 5) and sodium amidinate 57 to the exclusion of any mixed aggregate. DFT computations suggest that 57 is a disolvate.

$$\begin{array}{c|c}
N & C & N \\
\hline
N & NaHMDS \\
\hline
THF, 25 °C
\end{array}$$

$$\begin{array}{c}
N & Na \\
N & Na \\
N & SiMe_3
\end{array}$$

$$\begin{array}{c}
SiMe_3 \\
SiMe_3
\end{array}$$

$$\begin{array}{c}
57
\end{array}$$

Rate studies showed first-order decays and concentrationindependent values of  $k_{\rm obsd}$ , consistent with a first-order dependence on nitrile **6**. A plot of  $k_{\rm obsd}$  vs THF concentration

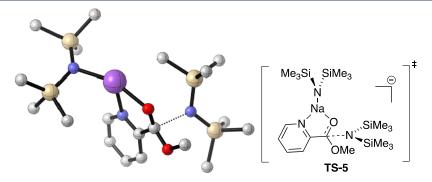
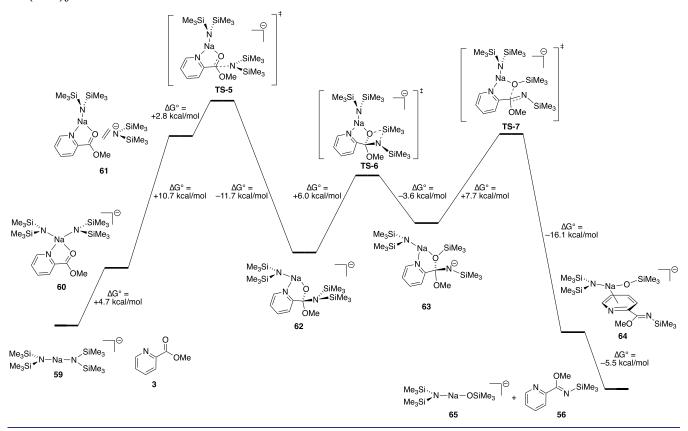
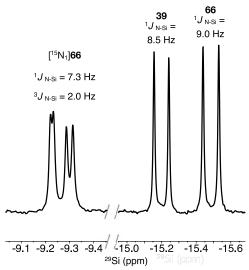


Figure 11. DFT-computed transition structure TS-5 for the NaHMDS-based addition to methyl-2-picolinate (3) solvated by THF. The implicit <sup>+</sup>Na(THF)<sub>6</sub> cation<sup>7b</sup> (Figure 12) is not included.

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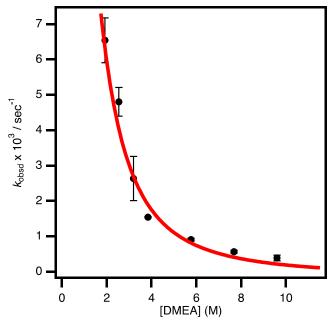
Scheme 6. DFT-Computed Reaction Coordinate for the Aminolysis of Methyl-2-Picolinate (3) with NaHMDS in THF with the <sup>+</sup>Na(THF)<sub>6</sub> Omitted for All Structures





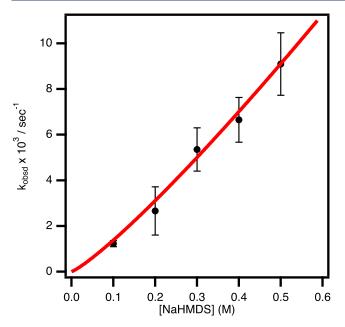
**Figure 13.** <sup>29</sup>Si NMR spectra of the [ $^{15}N_2$ ]NaHMDS dimer (39,  $^{1}J_{N-Si} = 8.5 \text{ Hz}$ ) and mixed aggregate [ $^{15}N_1$ ]66 ( $^{1}J_{N-Si} = 7.3 \text{ Hz}$ ,  $^{3}J_{N-Si} = 2.0 \text{ Hz}$ ).

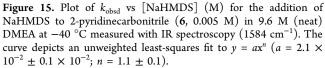
shows a distinct sigmoidal dependence affording a pronounced inhibition at high THF (Figure 16, curve A). Any doubt that this correlates with the NaHMDS dimer—monomer deaggregation (52 and 53, Scheme 5) was put to rest by superimposing the rate data on a plot of the equilibrium population of dimer (curve B) measured in 2020 by a different experimentalist. Clearly, deaggregation inhibits the reaction. A plot of  $k_{\rm obsd}$  vs NaHMDS in *neat* THF where NaHMDS is >98% monomer shows a clean second-order dependence

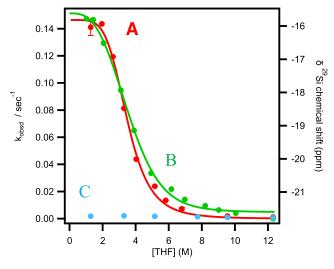


**Figure 14.** A plot of  $k_{\rm obsd}$  vs [DMEA] (M) in pentane for the addition of NaHMDS to pyridinecarbonitrile (6, 0.005 M) at -40 °C measured with IR spectroscopy (1584 cm<sup>-1</sup>). The curve depicts an unweighted least-squares fit to  $y = ax^n/(1+bx^n)$  ( $n = -1.9 \pm 0.1$ ;  $a = 2.3 \times 10^{-2} \pm 0.1 \times 10^{-2}$ ;  $b = 1.7 \times 10^{-1} \pm 0.1 \times 10^{-1}$ ).

(Figure 17), consistent with a requisite monomer-to-dimer aggregation before rate-limiting addition. The crudely defined zeroth-order THF dependence implicates the *partial* rate law in eq 9, pointing to the overall dimer-based transition

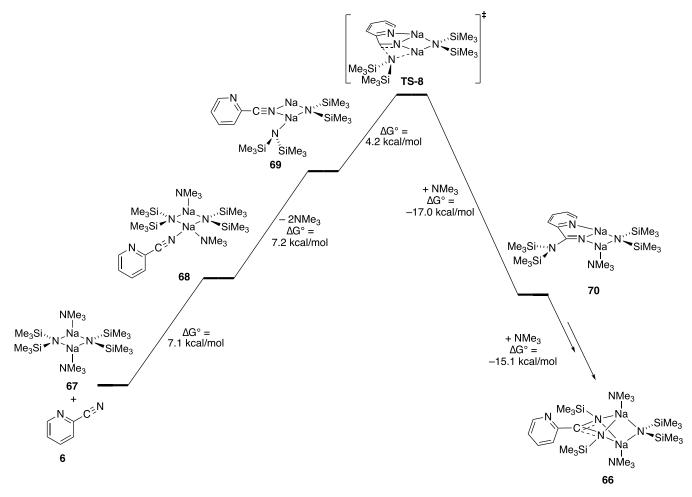


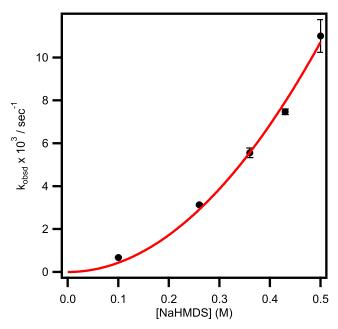




**Figure 16.** A plot of  $k_{\rm obsd}$  vs [THF] (M) for the addition of NaHMDS to 2-pyridinecarbonitrile in pentane cosolvent (curve A, red) and  $^{29}{\rm Si}$  chemical shift (curve B, green) plotted vs [THF] in 2:1 pentane/toluene as cosolvent measured at  $-20~{\rm ^{\circ}C}$ . The latter function fits to a model based on an  $A_2S_2-AS_4$  equilibrium (Supporting Information). Th, S2 The blue data (Curve C) derive from 2, S-Me<sub>2</sub>THF rather than pentane as cosolvent (vide infra).

## Scheme 7. DFT-Computed Reaction Coordinate for the Aminolysis of 2-Pyridinecarbonitrile (6) by NaHMDS Using $Me_3N$ as a DMEA Surrogate





**Figure 17.** Plot of  $k_{\rm obsd}$  vs [NaHMDS] (M) for the addition of NaHMDS to picolinonitrile (6, 0.005 M) in 12.8 M (neat) THF at  $-20~^{\circ}{\rm C}$  measured with IR spectroscopy (1584 cm<sup>-1</sup>). The curve depicts an unweighted least-squares fit to  $y = ax^n$  ( $a = 4.3 \times 10^{-2} \pm 0.1 \times 10^{-2}$ ;  $n = 2.0 \pm 0.1$ ).

structure,  $[A_2S_2(ArCN)]^{\frac{1}{4}.51,52}$  The DFT-computed transition structure **TS-10** is illustrated in Figure 18.

$$-d[\mathbf{6}]/\mathrm{d}t = k'[\mathbf{6}]^{1}[\mathrm{AS}_{4}]^{2} \tag{9}$$

The DFT-computed reaction coordinate for addition to nitrile 6 in THF shows many parallels with that in DMEA that are relegated to the Supporting Information. The notable differences are the exothermic solvation of all key species including transition structure TS-10 in Figure 18, mixed dimerbased transition structure TS-9 for silicon transfer, and unobserved mixed aggregate 71 whose analog (66, eq 7) was fully characterized in DMEA.<sup>53</sup>

We have accumulated evidence of preaggregation-based reactions originating from monomers, but none have ever been so poignant. Why is the dimer-based addition to nitriles dominant? A partial answer is that the trajectory of the silazide attack on the nitrile  $\pi$  system in **TS-10** (Figure 18) appears

optimal. The most stable transition structure for monomerbased addition (TS-11, Figure 19) is less favorable by 4.4 kcal/ mol using a monomer reference state, which appears to be a consequence of an inferior trajectory.

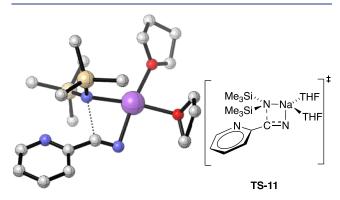


Figure 19. DFT-computed the transition structure TS-11 for the unfavorable NaHMDS-monomer-based addition to nitrile 6.

We would be remiss not to comment on the influence of 2,5-Me<sub>2</sub>THF rather than pentane as the cosolvent for nitrile addition (curve C in Figure 16). It provided further evidence of cooperative solvation. While checking for polarity effects, we detected and subsequently documented cooperative monomer solvation,  $(Me_3Si)_2NNa(THF)_x(Me_2THF)_y$ , noted in eq 6. The details are obscured in Figure 16 by an overlaying THF concentration dependence. The complete suppression of the rate suggested that the mixed solvated monomers persisted as an unproductive side equilibrium at stunningly low THF concentrations. Indeed, this was caused by a steep temperature dependence of the deaggregation. The message for us is that 2,5-Me<sub>2</sub>THF is a far better noncoordinating THF surrogate for lithium than for sodium.

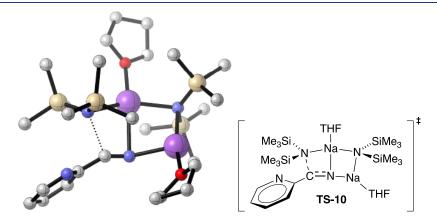


Figure 18. DFT-computed transition structure TS-10 for the NaHMDS-dimer-based addition to nitrile 6.

#### Scheme 8. Summary of NaHMDS Mechanistic Studies in THF Solution

Scheme 9. Summary of NaHMDS Mechanistic Studies in DMEA Solution

#### SUMMARY

The studies described above proved to be four discrete mechanistic studies in which the choice of solvent and conditions markedly influenced the outcome of the reactions. We offer an overview in which the results are grouped according to choice of THF or DMEA. This summary is organosodium centric; there is considerable organolsilicon chemistry (largely addressed computationally) that is left to the four parts described above. The arrows in Schemes 8 and 9 encompass many discrete steps including the organosilicon chemistry.

The THF results are summarized in Scheme 8. While NaHMDS in neat THF solution exists exclusively as tetrasolvated monomer 53, it reacts with methyl-2-picolinate, 3, via a dimer-based pathway in which triple-ion motif TS-5 is invoked at the rate-limiting transition structure. The critical chemoselectivity stems from a post-rate-limiting extrusion of NaOSiMe3 to form spectroscopically observable imino ether 56, which reacts further to give nitrile 6 and sodium amidinate 57 en route to a high-yielding one-pot synthesis of amidine 7. This ester-to-amidine conversion seems potentially important synthetically. Addition of NaHMDS to nitrile 6 is also dimerbased as depicted in computationally viable transition structure TS-10. The dominance of dimer-based reactivity under conditions affording monomeric NaHMDS should pique the interest of those interested in mechanistic alkali metal chemistry.

The chemistry of NaHMDS in poorly coordinating DMEA<sup>7b</sup> (Scheme 9) shows some parallels with the results in THF, but the differences are consequential. Although we do not consider a dimeric NaHMDS to be the proximate cause of dimer-based reactivity - many if not most monomer-based organoalkali metal reactions emanate from aggregates - the dominance of a dimer-based transition structure, TS-1, leading to observable mixed aggregate 43 aligns with previous studies of both LiHMDS and NaHMDS in DMEA.7c,54 In contrast to THF, however, a critical post-rate-limiting extrusion of MeOSiMe<sub>3</sub> rather than NaOSiMe<sub>3</sub> dictates the chemoselective formation of carboxamide 4. Observable mixed aggregate 50 is robust but finds a path to 66 over a month at 25 °C. Although this is clearly an inferior route to amidine 7 we examined the addition to nitrile 6 and, once again, uncovered dimer-based chemistry via an unsolvated transition structure, TS-8.

### CONCLUSIONS

While casting about for case studies to investigate solvent-structure—reactivity relationships in NaHMDS, we happened across several potentially useful reactions that are poorly represented in the literature and provided mechanistic insights that were wholly unexpected. The potential utility stems from C–N bond formation directly from aryl methyl esters, bypassing more activated forms. For us, however, they provided clean examples illustrating the role of solvation and aggregation in the chemistry of sodium amides in general and

NaHMDS in particular. The NaHMDS itself serves as a preface to ongoing studies of sodium alkyl(trimethylsilyl) amides with potentially greater utility as strong bases and for C–N bond-forming reactions.

The dimer- and mixed dimer-based reactivity has precedent, 7c,36 but the total dominance of aggregate-based reactivity still came as a surprise. From a mechanistically tactical perspective, the work underscores both the power of <sup>29</sup>Si-<sup>15</sup>N coupling to determine solution structures and the general merits of <sup>29</sup>Si NMR spectroscopy. (We asserted previously that <sup>29</sup>Si NMR spectroscopy is underutilized by those outside the organosilicon community.) We offer a final caveat to those who might be tempted to probe primary- vs secondary-shell solvation by using 2,5-Me<sub>2</sub>THF as a polar, noncoordinating cosolvent. Despite its success in organolithium chemistry, the larger sodium ion may call for the more expensive 2,2,5,5-Me<sub>4</sub>THF. On the truism that an effect is either sterics or electronics, ours' and others' experience with metal ion solvation is that sterics dominates.<sup>55</sup>

#### EXPERIMENTAL SECTION

**Reagents and Solvents.** NaHMDS and [<sup>15</sup>N]NaHMDS were prepared as white crystalline solids. Toluene, hexanes, THF, MTBE, cyclopentane, 2,5-Me<sub>2</sub>THF, and HMPA were distilled from blue or purple solutions containing sodium benzophenone ketyl. All substrates and products in Table 1 are commercially available. <sup>31,32</sup>

General Procedure A: Picolinamide 4. Solid sodium hexamethyldisilazide (NaHMDS, 165 mg, 0.9 mmol) was dissolved in 2.0 mL of DMEA at 25  $^{\circ}\text{C}.$  Two mL of the NaHMDS solution was added to a dry 5 mL Kimble vial equipped with a magnetic stir bar. Methyl-2-picolinate (3, 36  $\mu$ L, 0.30 mmol) was then added to the reaction solution. The reaction mixture was stirred at 25 °C for 0.3 h. DI water (1 mL) was added, and the resulting biphasic mixture was partitioned between water (1 mL) and ethyl ether (2 mL). The aqueous layer was separated and extracted further with three 2 mL portions of ethyl ether. The combined organic layers were dried over anhydrous magnesium sulfate and then concentrated. Purification of the residue by flash column chromatography (80% ethyl acetate in hexanes) afforded picolinamide 4 as a white solid. <sup>1</sup>H NMR (500 MHz, CDCl3)  $\delta$  8.56 (d, J = 4.6 Hz, 1H), 8.20 (d, J = 7.8 Hz, 1H), 7.89 (s, 1H), 7.84 (td, J = 7.7, 1.8 Hz, 1H), 7.43 (dd, J = 7.6, 4.8 Hz, 1H), 6.23 (s, 1H).  ${}^{13}C\{{}^{1}H\}$  NMR (126 MHz, CDCl<sub>3</sub>)  $\delta$ : 166.99, 145.59, 148.32, 137.31, 126.47, 122.44. HRMS (DART) Calc. for  $C_6H_6N_2O$  (M + H<sup>+</sup>): 123.04746; found: 123.05493.

General Procedure B: Amidine 7. Solid sodium hexamethyldisilazide (NaHMDS, 165 mg, 0.9 mmol) was dissolved in 1.5 mL of THF at 25 °C. 1.5 mL of the NaHMDS solution was added to a dry 5 mL Kimble vial equipped with a magnetic stir bar. Methyl picolinate (3, 36  $\mu$ L, 0.30 mmol) was then added to the reaction solution. The reaction mixture was stirred at 50 °C for 0.3 h. DI water (1 mL) was then added. 2 M HCl aqueous solution was added to adjust pH = 1, which then was stirred at 25 °C for 1 h. Saturated NaOH solution was added until pH = 12, and the mixture was partitioned between water (1 mL) and chloroform (4 mL). The aqueous layer was separated and extracted further with three 4 mL portions of chloroform. The combined organic layers were dried over anhydrous magnesium sulfate and then concentrated to afford amidine 7 as a yellow oil. <sup>1</sup>H NMR (500 MHz, CDCl3)  $\delta$  8.56 (m, 1H), 8.11 (d, J = 7.9 Hz, 1H), 7.78 (td, I = 7.7, 1.8 Hz, 1H), 7.35 (dd, I = 7.5, 4.9 Hz, 1H), 5.94 (s, 3H).  $^{13}$ C $\{^{1}$ H $\}$  NMR (126 MHz, CDCl<sub>3</sub>)  $\delta$ : 161.06, 150.79, 148.35, 137.03, 125.21, 120.81. HRMS (DART) Calc. for  $C_6H_7N_3$  (M + H<sup>+</sup>): 122.06345; found: 122.07097.

**NMR Spectroscopic Analyses.** An NMR tube fitted with a double-septum under vacuum was flame-dried on a Schlenk line and allowed to passively cool to room temperature, backfilled with argon, and placed in a dry ice/acetone cooling bath. Individual stock solutions of the substrate and NaHMDS or [<sup>15</sup>N]NaHMDS were

prepared at room temperature. The appropriate amounts of the substrate, NaHMDS, solvent, and cosolvent were added sequentially to the tube cooled to  $-78~^{\circ}\mathrm{C}$  via a gastight syringe. The tube was flame-sealed under a partial vacuum while cold to minimize evaporation in some cases and left unsealed for incremental additions. The tubes were mixed with a vortex mixer for approximately 10 s to minimize warming. Standard  $^{1}\mathrm{H},~^{13}\mathrm{C},$  and  $^{29}\mathrm{Si}$  direct detection spectra were recorded at 500, 125.79, and 99.36 MHz, respectively, and referenced to Me<sub>4</sub>Si (0.0 ppm). Integration of the NMR signals was determined using the line-fitting method included in MNova (Mestrelab research S.L.).

Rate Studies. IR spectra were recorded with an in situ IR spectrometer fitted with a 30-bounce, silicon-tipped probe. The spectra were acquired at a gain of 1 and a resolution of 4 cm<sup>-1</sup>. All tracked reactions were conducted under a positive flow of argon from a Schlenk line. A representative reaction was carried out as follows: The IR probe was inserted through a Teflon adapter and an O-ring seal into an oven-dried, cylindrical flask fitted with a magnetic stir bar and a T-joint. The T-joint was capped with a septum for injections and an argon line. After evacuation under full vacuum, heating, and flushing with argon, the flask was charged with the solvent mixtures of our choice (toluene, DMEA, THF, 2,5-Me<sub>2</sub>THF, and 2,2,5,5-Me<sub>4</sub>THF) and cooled to -78 °C in a dry ice-acetone bath. A set of 256 baseline scans were collected, and IR spectra were recorded every 15 s from 30 scans. The reaction vessel was charged with stock solutions of NaHMDS and additional cosolvent through the septum. One set of scans was collected before the addition of substrate through the septum, and the baseline is zeroed. The substrate was added neat or in highly concentrated solutions and was tracked to completion (1731 cm<sup>-1</sup> for 7, for example). The spectrometer was configured to collect spectra every 5 s from 16 scans. The reaction was tracked over 3-5 half-lives monitoring the disappearance of the starting material and the appearance of the product. The former was used for the rate studies.

**Density Functional Theory (DFT) Computations.** All DFT calculations were carried out using Gaussian 16.<sup>29</sup> Prompted by a recent benchmarking of modern density functionals, all calculations were conducted at the M06-2X level of theory.<sup>30a-c</sup> A pruned (99,590) integration grid (equivalent to Gaussian's "UltraFine" option) was used for all calculations. The Ahlrichs basis set def2-svp was used for geometry optimizations and the expanded def2-tsvp basis set for single-point energy calculations.<sup>30d</sup> Ball-and-stick models were rendered using CYLview 1.0b.<sup>30e</sup> A large number of DFT-computed energies are archived in the Supporting Information.

#### ASSOCIATED CONTENT

#### **Supporting Information**

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/jacs.3c07317.

Synthetic and experimental procedures, <sup>1</sup>H and <sup>29</sup>Si NMR spectroscopic, rate, and computational data (PDF)

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#### Notes

The authors declare no competing financial interest.

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- (52) The order in THF cannot be determined from a fit with values of -4 through -6 all providing adequate fits to the data. The presumed value of -6 comes from the assigned solvation states of NaHMDS monomer as tetrasolvated,  $^{7b}$  and transition structure **50** as a disolvate.
- (53) A monomer-based N-to-N' silicon transfer analogous to that depicted in Figure 18 is substantially less favorable.
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